

Modified Canny Edge Detection Algorithm

Notations:

X_{raw} = raw RGB image

$$X_{raw} \in \mathbb{R}^{3 \times M \times N}$$

where M = number of rows, N = number of columns

X_{raw} = raw pixel value of c^{th} color channel, i^{th} row, j^{th} column

$$edge_list \in \mathbb{R}^{num_edges \times 2 \times 2}$$

$edge_list[e, v, i]$ = value of e^{th} edge, v^{th} vertex, i^{th} index

All arrays (matrices and tensors) are indexed from zero in this document.

1. Step 1: Gaussian Blur Filter:

1.1. Inputs:

1.1.1. Symmetric 2D Gaussian kernel matrix: $G \in \mathbb{R}^{K \times K}$

1.1.1.1. Sample symmetric 2D Gaussian distribution:

$$G[x + p, y + p] = \frac{1}{2\pi\sigma^2} \exp(-(x^2 + y^2)/(2\sigma^2)), \quad \text{for } x, y \in \{-p, \dots, p\},$$

$$\text{where } p = (K - 1)/2$$

1.1.2. Normalize kernel (so that its elements sum to 1):

$$G[i, j] := G[i, j] / \sum_{i=0}^{K-1} \sum_{j=0}^{K-1} G[i, j], \quad \text{for } i, j \in \{0, \dots, K - 1\}$$

1.1.3. Zero-padded raw image: $\tilde{X}_{raw} \in \mathbb{R}^{3 \times (M+2p) \times (N+2p)}$

1.1.3.1. Zero-pad raw image for 2D “same” convolution:

$$\tilde{X}_{raw}[c, i, j] = \begin{cases} 0, & i < p \text{ or } j < p \text{ or } i > M - 1 + p \text{ or } j > N - 1 + p \\ X_{raw}[c, i - p, j - p], & \text{otherwise} \end{cases},$$

$$\text{for } c \in \{0, 1, 2\}, i \in \{0, \dots, (M + 2p) - 1\}, j \in \{0, \dots, (N + 2p) - 1\},$$

$$\text{where } p = (K - 1)/2$$

1.2. Computation:

1.2.1. Compute 2D “same” convolution:

$$X_{blur}[c, i, j] = \sum_{m=i}^{i+K-1} \sum_{n=j}^{j+K-1} \tilde{X}_{raw}[c, m, n] * G[m - i, n - j],$$

$$\text{for } c \in \{0, 1, 2\}, i \in \{0, \dots, M - 1\}, j \in \{0, \dots, N - 1\}$$

1.3. Outputs:

1.3.1. Blurred image: $X_{blur} \in \mathbb{R}^{3 \times M \times N}$

2. Step 2: Gradient Estimation:

2.1. Inputs:

2.1.1. Sobel operator kernels:

2.1.1.1. Horizontal Sobel operator (for horizontal gradients):

$$S_{horiz} = \frac{1}{8} \begin{bmatrix} -1 & 0 & +1 \\ -2 & 0 & +2 \\ -1 & 0 & +1 \end{bmatrix}$$

2.1.1.2. Vertical Sobel operator (for vertical gradients):

$$S_{vert} = \frac{1}{8} \begin{bmatrix} +1 & +2 & +1 \\ 0 & 0 & 0 \\ -1 & -2 & -1 \end{bmatrix}$$

2.1.2. Blurred image: $X_{blur} \in \mathbb{R}^{3 \times M \times N}$

2.2. Computation:

2.2.1. Convolve with horizontal and vertical Sobel operators (post zero-padded):

$$W_d[c, i, j] = \begin{cases} 0, & i = 0 \text{ or } j = 0 \text{ or } i = M - 1 \text{ or } j = N - 1 \\ \sum_{m=i-1}^{i+1} \sum_{n=j-1}^{j+1} X_{blur}[c, m, n] * S_d[m - (i - 1), n - (j - 1)], & \text{otherwise} \end{cases}$$

for $d \in \{horiz, vert\}, c \in \{0, 1, 2\}, i \in \{0, \dots, M - 1\}, j \in \{0, \dots, N - 1\}$

2.2.2. Compute L1-norm of horizontal and vertical gradients:

$$W[c, i, j] = |W_{horiz}[c, i, j]| + |W_{vert}[c, i, j]|,$$

for $c \in \{0, 1, 2\}, i \in \{0, \dots, M - 1\}, j \in \{0, \dots, N - 1\}$

2.2.3. Take maximum gradient over color channels:

$$Y[i, j] = \max_{c \in \{0, 1, 2\}} W[c, i, j], \quad \text{for } i \in \{0, \dots, M - 1\}, j \in \{0, \dots, N - 1\}$$

2.3. Outputs:

2.3.1. Gradient image: $Y \in \mathbb{R}^{M \times N}$

3. Step 3: Horizontal Non-Maximal Suppression:

3.1. Inputs:

3.1.1. Gradient Image: $Y \in \mathbb{R}^{M \times N}$

3.1.2. (Half) suppression length: r

3.2. ²Algorithm:

```
for  $i = 0$  to  $M - 1$ :  
  for  $j = 0$  to  $N - 1$ :  
    if  $j \leq r - 1$ :  
      if  $Y[i, j] \neq \max_{j \leq k \leq j+r} Y[i, k]$ :  
         $Y[i, j] := 0$   
    else if  $j \geq N - r$ :  
      if  $Y[i, j] \neq \max_{j-r \leq k \leq j} Y[i, k]$ :  
         $Y[i, j] := 0$   
    else:  
      if  $Y[i, j] \neq \max_{j-r \leq k \leq j+r} Y[i, k]$ :  
         $Y[i, j] := 0$ 
```

3.3. Outputs:

3.3.1. Non-maximally suppressed gradient image: $Y \in \mathbb{R}^{M \times N}$

4. Step 4: Long Vertical Edge Determination:

4.1. Inputs:

- 4.1.1. Non-maximally suppressed gradient image: $Y \in \mathbb{R}^{M \times N}$
- 4.1.2. Low and high thresholds: $thresh_{low}, thresh_{high}$
- 4.1.3. Vertical and horizontal “scanning” lengths: $scan_{vert}, scan_{horiz}$
- 4.1.4. Minimum edge length: min_edge_length

4.2. Algorithm (Simplified):

```
num_edges = 0
for each pixel  $Y[i, j]$  in  $Y$ :
    if  $Y[i, j] \geq thresh_{high}$ :
        start =  $[i, j]$ 
        connect = true
        current_pix = start
        while connect == true and current_pix with scan lengths fits in  $Y$ :
             $max_{value} = \max_{i+1 \leq m \leq i+scan_{vert}, j-scan_{horiz} \leq n \leq j+scan_{horiz}} Y[m, n]$ 
             $max_{ind} = \arg \max_{i+1 \leq m \leq i+scan_{vert}, j-scan_{horiz} \leq n \leq j+scan_{horiz}} Y[m, n]$ 
            if  $max_{value} \geq thresh_{low}$ :
                current_pix =  $max_{ind}$ 
            else:
                connect = False
        stop = current_pix
        if  $|start[0] - stop[0]| + |start[1] - stop[1]| \geq min\_edge\_length$ :
            edge_list[num_edges] = [start, stop]
            num_edges += 1
```

4.3. Outputs:

- 4.3.1. Array of edges: $edge_list \in \mathbb{R}^{num_edges \times 2 \times 2}$

¹ In step 1.1.1 and 1.1.2, two different sets of indices are used: (x, y) and (i, j) . The index set (x, y) is used in step 1.1.1 because it is more natural for a 2D Gaussian distribution. In addition, (x, y) is used in order to avoid possible confusion of i or j being interpreted as the imaginary unit.

² In step 3.2, pseudocode is used instead of a mathematical notation (as in steps 1 and 2), because it is more understandable in pseudocode in this particular case.